

RAW SEQUENCE LISTING
PATENT APPLICATION US/08/653,114DATE: 08/07/96
TIME: 10:38:04 #16

INPUT SET: S2729.raw

This Raw Listing contains the General
Information Section and up to the first 5 pages.

SEQUENCE LISTING

ENTERED

(1) General Information:

- (i) APPLICANT: Falck-Pedersen, Erik S.
- (ii) TITLE OF INVENTION: ADENOVIRUS GENE EXPRESSION SYSTEM
- (iii) NUMBER OF SEQUENCES: 1
- (iv) CORRESPONDENCE ADDRESS:
 - (A) ADDRESSEE: Alan S. Korman
 - (B) STREET: 1600 Main Place Tower
 - (C) CITY: Buffalo
 - (D) STATE: New York
 - (E) COUNTRY: U.S.A.
 - (F) ZIP: 14202
- (v) COMPUTER READABLE FORM:
 - (A) MEDIUM TYPE: Floppy disk
 - (B) COMPUTER: IBM PC compatible
 - (C) OPERATING SYSTEM: PC-DOS/MS-DOS
 - (D) SOFTWARE: PatentIn Release #1.0, Version #1.25
- (vi) CURRENT APPLICATION DATA:
 - (A) APPLICATION NUMBER: US/08/653,114
 - (B) FILING DATE:
 - (C) CLASSIFICATION:
- (vii) PRIOR APPLICATION DATA:
 - (A) APPLICATION NUMBER: 08/166,925
 - (B) FILING DATE: 12/14/93
- (viii) ATTORNEY/AGENT INFORMATION:
 - (A) NAME: Korman, Alan S.
 - (B) REGISTRATION NUMBER: 33,932
 - (C) REFERENCE/DOCKET NUMBER: 19603/230
- (ix) TELECOMMUNICATION INFORMATION:
 - (A) TELEPHONE: 716-853-8104
 - (B) TELEFAX: 716-853-8109

(2) INFORMATION FOR SEQ ID NO:1:

- (i) SEQUENCE CHARACTERISTICS:

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TIME: 10:38:10

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47 (A) LENGTH: 6783 base pairs
48 (B) TYPE: nucleic acid
49 (C) STRANDEDNESS: double
50 (D) TOPOLOGY: linear
51
52 (ii) MOLECULE TYPE: cDNA (genomic)
53
54
55
56 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1:
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58 TTCCATCATC AATAATATAC CTTATTTTGG ATTGAAGCCA ATATGATAAT GAGGGGGTGG 60
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62 TGTGATGTTG CAAGTGTGGC GGAACACATG TAAGCGACGG ATGTGGCAAA AGTGACGTTT 180
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98 TTGAGTGACA ATGACATCCA CTTTGCCTTT CTCTCCACAG GTGTCCACTC CCAGGTCCAA 1260
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203	TTGATGCGTT TCTTACCTCT GGTTC CATG AGCCGGTGTC CACGCTCGGT GACGAAAAGG	4440
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212	GACCGCTTTC	GCTGGAGCGC	GACGATGATC	GGCCTGTTCG	TTGCGGTATT	CGGAATCTTG	4680
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214	CACGCCCTCG	CTCAAGCCTT	CGTCACTGGT	CCCCCCACCA	AACGTTTCGG	CGAGAAGCAG	4740
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228	ACCGGATACC	TGTCCGCCTT	TCTCCCTTCG	GGAAGCGTGG	CGCTTTCTCA	ATGCTCACGC	5160
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230	TGTAGGTATC	TCAGTTCGGT	GTAGGTCGTT	CGCTCCAAGC	TGGGCTGTGT	GCACGAACCC	5220
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242	ACGCGCAGAA	AAAAAGGATC	TCAAGAAGAT	CCTTTGATCT	TTTCTACGGG	GTCTGACGCT	5580
243							
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258	AATAGTTTGC	GCAACGTTGT	TGCCATTGCT	GCAGGCATCG	TGGTGTACAG	CTCGTCGTTT	6060

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SEQUENCE VERIFICATION REPORT
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